## Biocellion: A Large Capacity Modeling Platform for Multicellular Biological Systems Seunghwa Kang, Pacific Northwest National Laboratory & Northwest Institute for Advanced Computing Simon Kahan, Biocellion Social Purpose Corporation

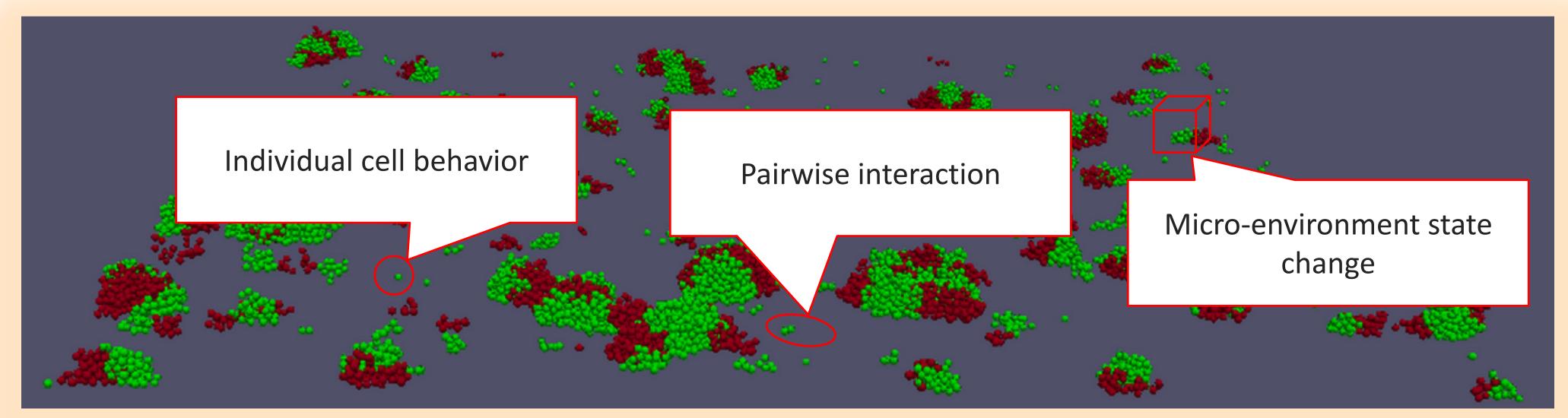


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Studying emergent many-cell behaviors demands unprecedented capacity: high-performance simulation of flexible models at enormous scale.

Modeling Capacity = Performance (X) Flexibility

## Attention to basic building blocks common to modeling all living systems...

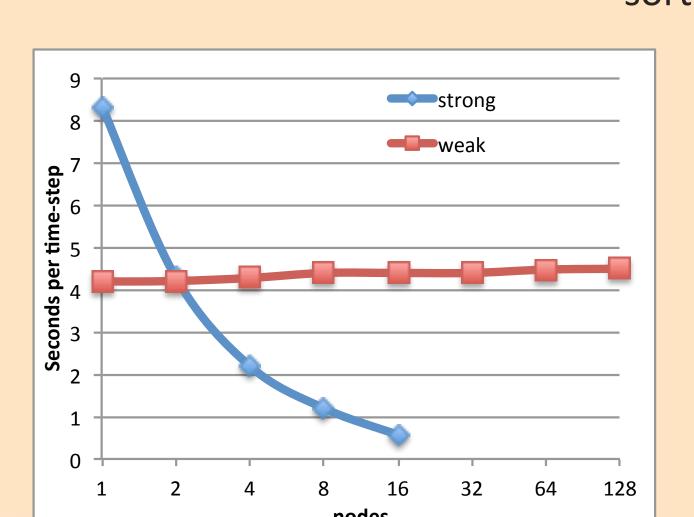


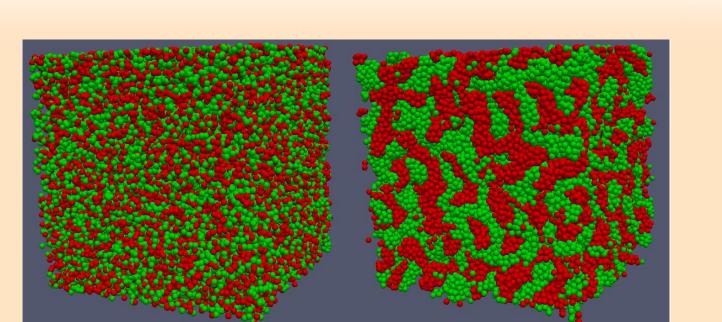
Self: cell cycle, geometry, growth, division, migration, ... ODEs within each cell

**Neighbor interactions:** shoving, adhesion, phagocytosis, anastomosis, ... short range "N-body" problems **Long-range interactions:** secretion, uptake, diffusion, advection, ... PDEs over whole domain

## ... that leverages our investment in performance-tuning simulation software,

"Cell sorting" occurs when cells adhere preferentially to cells of the same type, forming clumps. Cell sorting serves as a synthetic benchmark for simulation performance. The literature reports no results sorting more than 2M cells. Biocellion sorts >1B cells.





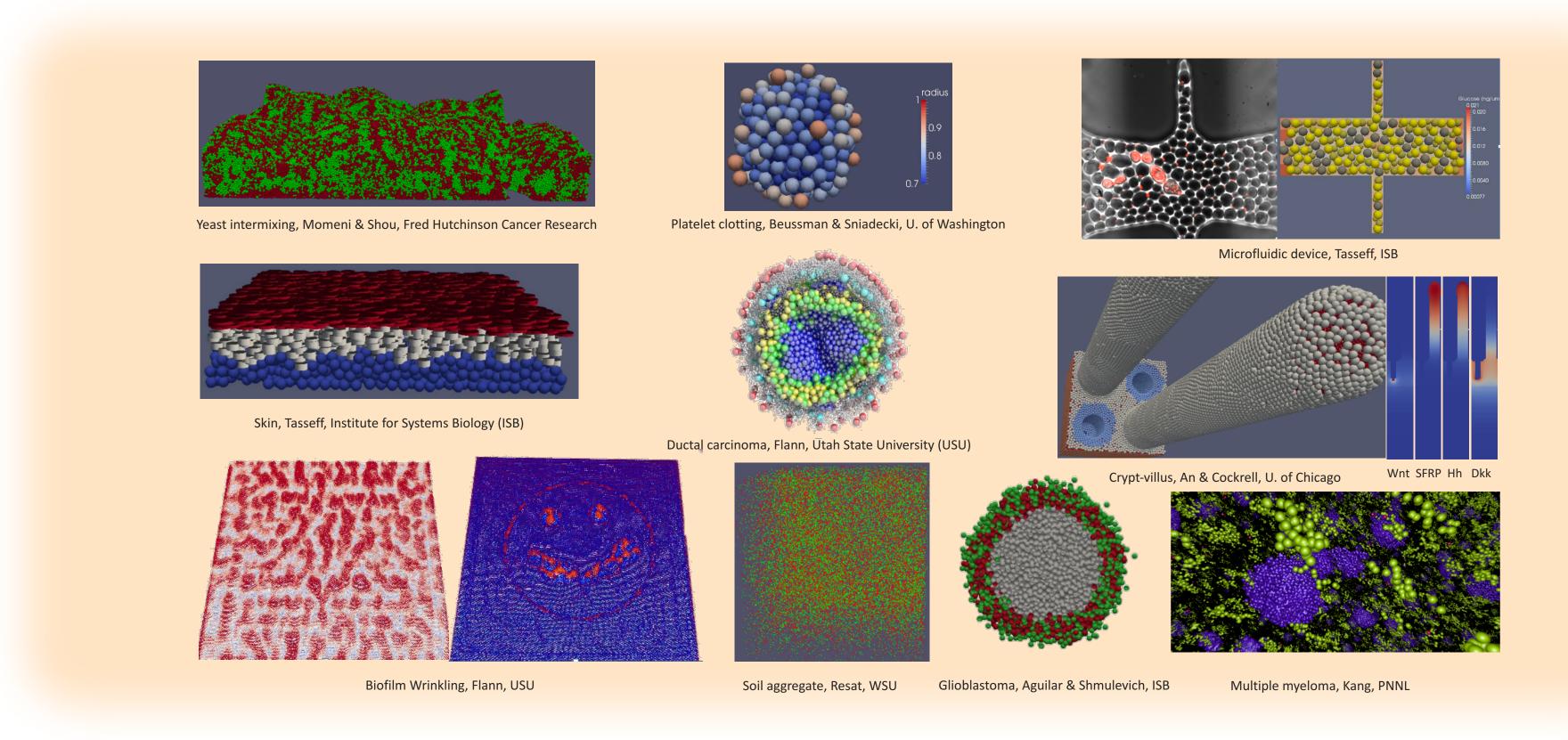
Randomly distributed modeled cells (left) migrate to form the clumped patterns (right).

Biocellion sorted 26.8M cells almost 16 times as quickly with 16 computing nodes as with 1 (blue curve): 400x faster than any published result! The number of cells that can be sorted in any fixed amount of time is proportional to the size of computer used (red): on 128 nodes, 128x13.4M = 1.7B cells are sorted almost as fast as 13.4M on 1.

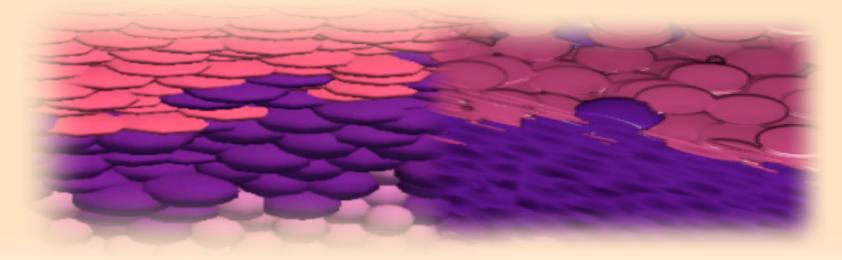
in extensibility,

and in accommodating many-cell models.

## ... yields the flexibility needed to model in many domains...



If you cannot incrementally refine your model, you risk wasting your investment. Large modeling capacity allows you to incrementally adopt advances in biological system modeling. For example, a sphere is a good first approximation of most cell geometries. A second approximation better capturing mechanical interactions responsible for tissue morphology is the ellipsoid. Biocellion supports interaction mechanics of arbitrarily oriented ellipsoidal cells:



When initializing cell placement in a virtual experiment, can you extract billions of cell locations from 3D reconstruction of histology slices? Not likely. Biocellion provides built-in support for packing cells along surfaces or between surfaces defined by  $f_i(x,y)$  you provide.



So your investment in modeling simulates on laptop to supercomputer with no added work.



See other Biocellion modeling posters here:: Skin (55), Biofilm Morphology (67), Multiple Myeloma (79)





All you need to get started...

**Prerequisites:** 

C & Linux programming skills Mathematical modeling skills

Domain measurements, knowledge and calibration source

**Download**: biocellion.com/download **Manuals**: biocellion.com/documentation

**Tutorial**: biocellion.com/wiki\_root/index.php/Getting\_Started **Support**: groups.google.com/forum/#!forum/biocellion-support



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